



# SEQUENCE LISTING

<110> Chuntharapai, Anan  
Grewal, Iqbal  
Kim, Kyung Jin  
Yan, Minhong

<120> TACI Antibodies and Uses Thereof

<130> 50474/017002

<140> US 10/626,914

<141> 2003-07-25

<150> US 60/398,530

<151> 2002-07-25

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Ser Cys Lys Thr Ile Cys Asn His Gln Ser Gln Arg Thr Cys Ala Ala  
 50 55 60

Phe Cys Arg Ser Leu Ser Cys Arg Lys Glu Gln Gly Lys Phe Tyr Asp  
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His Leu Leu Arg Asp Cys Ile Ser Cys Ala Ser Ile Cys Gly Gln His  
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Pro Lys Gln Cys Ala Tyr Phe Cys Glu Asn Lys Leu Arg Ser Pro Val  
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Cys Ser Cys Gln Pro Arg Ser Arg Pro Arg Gln Ser Pro Ala Lys Ser	195	200	205
Ser Gln Asp His Ala Met Glu Ala Gly Ser Pro Val Ser Thr Ser Pro	210	215	220
Glu Pro Val Glu Thr Cys Ser Phe Cys Phe Pro Glu Cys Arg Ala Pro	225	230	235
Thr Gln Glu Ser Ala Val Thr Pro Gly Thr Pro Asp Pro Thr Cys Ala	245	250	255
Gly Arg Trp Gly Cys His Thr Arg Thr Thr Val Leu Gln Pro Cys Pro	260	265	270
His Ile Pro Asp Ser Gly Leu Gly Ile Val Cys Val Pro Ala Gln Glu	275	280	285
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Pro Pro Leu Thr Cys Gln Arg Tyr Cys Asn Ala Ser Val Thr Asn Ser  
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Val Lys Gly Thr Asn Ala Ile Leu Trp Thr Cys Leu Gly Leu Ser Leu  
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Ile Ile Ser Leu Ala Val Phe Val Leu Met Phe Leu Leu Arg Lys Ile  
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Ser Ser Glu Pro Leu Lys Asp Glu Phe Lys Asn Thr Gly Ser Gly Leu  
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Leu Gly Met Ala Asn Ile Asp Leu Glu Lys Ser Arg Thr Gly Asp Glu  
100 105 110

Ile Ile Leu Pro Arg Gly Leu Glu Tyr Thr Val Glu Glu Cys Thr Cys  
115 120 125

Glu Asp Cys Ile Lys Ser Lys Pro Lys Val Asp Ser Asp His Cys Phe  
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Pro Leu Pro Ala Met Glu Glu Gly Ala Thr Ile Leu Val Thr Thr Lys  
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Ala Ala Thr Leu Leu Leu Ala Leu Leu Ser Cys Cys Leu Thr Val Val  
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Ser Phe Tyr Gln Val Ala Ala Leu Gln Gly Asp Leu Ala Ser Leu Arg  
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Ala Glu Leu Gln Gly His His Ala Glu Lys Leu Pro Ala Gly Ala Gly  
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Ala Pro Lys Ala Gly Leu Glu Glu Ala Pro Ala Val Thr Ala Gly Leu  
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Lys Ile Phe Glu Pro Pro Ala Pro Gly Glu Gly Asn Ser Ser Gln Asn  
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Ser Arg Asn Lys Arg Ala Val Gln Gly Pro Glu Glu Thr Val Thr Gln  
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Asp Cys Leu Gln Leu Ile Ala Asp Ser Glu Thr Pro Thr Ile Gln Lys  
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Gly Ser Tyr Thr Phe Val Pro Trp Leu Leu Ser Phe Lys Arg Gly Ser  
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Gly His Leu Ile Gln Arg Lys Lys Val His Val Phe Gly Asp Glu Leu  
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Ser Leu Val Thr Leu Phe Arg Cys Ile Gln Asn Met Pro Glu Thr Leu  
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Pro Asn Asn Ser Cys Tyr Ser Ala Gly Ile Ala Lys Leu Glu Glu Gly  
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Gly Glu Arg Ser Arg Lys Arg Arg Ala Val Leu Thr Gln Lys Gln Lys  
100 105 110

Lys Gln His Ser Val Leu His Leu Val Pro Ile Asn Ala Thr Ser Lys  
115 120 125

Asp Asp Ser Asp Val Thr Glu Val Met Trp Gln Pro Ala Leu Arg Arg  
130 135 140

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145 150 155 160

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Thr Met Gly Gln Val Val Ser Arg Glu Gly Gln Gly Arg Gln Glu Thr  
180 185 190

Leu Phe Arg Cys Ile Arg Ser Met Pro Ser His Pro Asp Arg Ala Tyr  
195 200 205

Asn Ser Cys Tyr Ser Ala Gly Val Phe His Leu His Gln Gly Asp Ile  
210 215 220

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<212> DNA

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cctgagggac tgcatacagct gtgcctccat ctgtggacag caccctaagc aatgtgcata 180  
cttctgtgag aacaagctca ggagcccagt gaaccttcca ccagagctca ggagacagcg 240  
gagtggagaa gttgaaaaca attcagacaa ctcggaagag taccaaggat tggagcacag 300  
aggctcagaa gcaagtccag ctctcccggg gctgaagctg agtgcagatc aggtggccct 360  
ggctctacagc acgctggggc tctgcctgtg tgccgtcctc tgctgcttcc tgggtggcgg 420  
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ggagagcgca gtcacgcctg ggacccccga cccacttgt gctggaaggt gggggtgcca 660  
caccaggacc acagtccctgc agccttgccc acacatccca gacagtggcc ttggcattgt 720  
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ggagggagag agatggagag gaggggagag agaaagagag gtggggagag gggagagaga 840  
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gagagagaca gagggagaga gagacagagg ggaagagagg cagagaggga aagaggcaga 960  
gaaggaaaaga gacaggcaga gaaggagaga ggcagagagg gagagaggca gagagggaga 1020  
gaggcagaga gacagagagg gagagaggga cagagagaga tagagcagga ggtcggggca 1080  
ctctgagtcc cagttcccag tgcagctgta ggtcgtcatc acctaaccac acgtgcaata 1140  
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ttggcagctg cccttctca aaaaaaaaaa aaaaaaaaaa 1239

<210> 14  
<211> 247  
<212> PRT  
<213> Homo sapiens

<400> 14

Met Ser Gly Leu Gly Arg Ser Arg Arg Gly Gly Arg Ser Arg Val Asp  
1 5 10 15

Gln Glu Glu Arg Trp Ser Leu Ser Cys Arg Lys Glu Gln Gly Lys Phe  
20 25 30

Tyr Asp His Leu Leu Arg Asp Cys Ile Ser Cys Ala Ser Ile Cys Gly  
 35 40 45

Gln His Pro Lys Gln Cys Ala Tyr Phe Cys Glu Asn Lys Leu Arg Ser  
 50 55 60

Pro Val Asn Leu Pro Pro Glu Leu Arg Arg Gln Arg Ser Gly Glu Val  
 65 70 75 80

Glu Asn Asn Ser Asp Asn Ser Gly Arg Tyr Gln Gly Leu Glu His Arg  
 85 90 95

Gly Ser Glu Ala Ser Pro Ala Leu Pro Gly Leu Lys Leu Ser Ala Asp  
 100 105 110

Gln Val Ala Leu Val Tyr Ser Thr Leu Gly Leu Cys Leu Cys Ala Val  
 115 120 125

Leu Cys Cys Phe Leu Val Ala Val Ala Cys Phe Leu Lys Lys Arg Gly  
 130 135 140

Asp Pro Cys Ser Cys Gln Pro Arg Ser Arg Pro Arg Gln Ser Pro Ala  
 145 150 155 160

Lys Ser Ser Gln Asp His Ala Met Glu Ala Gly Ser Pro Val Ser Thr  
 165 170 175

Ser Pro Glu Pro Val Glu Thr Cys Ser Phe Cys Phe Pro Glu Cys Arg  
 180 185 190

Ala Pro Thr Gln Glu Ser Ala Val Thr Pro Gly Thr Pro Asp Pro Thr  
 195 200 205

Cys Ala Gly Arg Trp Gly Cys His Thr Arg Thr Thr Val Leu Gln Pro  
 210 215 220

Cys Pro His Ile Pro Asp Ser Gly Leu Gly Ile Val Cys Val Pro Ala  
 225 230 235 240

Gln Glu Gly Gly Pro Gly Ala  
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<210> 15

<211> 595  
 <212> DNA  
 <213> Homo sapiens

<400> 15  
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 cctgcgcacg ccgcggccga aaccggccgg ggccagcagc cctgcgcca ggacggcgct 180  
 gcagccgcag gaggcggtag gcgcgggggc cggcgaggcg gcgctgcccc tgccccgggt 240  
 gctctttggc gccccgcgc tgctgggcct ggcaactggc ctggcgctgg tctggtggg 300  
 tctggtgagc tggaggcggc gacagcggcg gcttcgcggc gcgtcctccg cagaggcccc 360  
 cgacggagac aaggacgccc cagagcccct ggacaaggc atcattctgt ctccgggaat 420  
 ctctgatgcc acagctcctg cctggcctcc tctggggaa gaccaggaa ccacccacc 480  
 tggccacagt gtccctgtgc cagccacaga gctgggctcc actgaactgg tgaccacaa 540  
 gacggccggc cctgagcaac aatagcaggg agccggcagg aggtggcccc tgccc 595

<210> 16  
 <211> 184  
 <212> PRT  
 <213> Homo sapiens

<400> 16  
 Met Arg Arg Gly Pro Arg Ser Leu Arg Gly Arg Asp Ala Pro Ala Pro  
 1 5 10 15  
 Thr Pro Cys Val Pro Ala Glu Cys Phe Asp Leu Leu Val Arg His Cys  
 20 25 30  
 Val Ala Cys Gly Leu Leu Arg Thr Pro Arg Pro Lys Pro Ala Gly Ala  
 35 40 45  
 Ser Ser Pro Ala Pro Arg Thr Ala Leu Gln Pro Gln Glu Ser Val Gly  
 50 55 60  
 Ala Gly Ala Gly Glu Ala Ala Leu Pro Leu Pro Gly Leu Leu Phe Gly  
 65 70 75 80  
 Ala Pro Ala Leu Leu Gly Leu Ala Leu Val Leu Ala Leu Val Leu Val  
 85 90 95

Gly Leu Val Ser Trp Arg Arg Arg Gln Arg Arg Leu Arg Gly Ala Ser  
100 105 110

Ser Ala Glu Ala Pro Asp Gly Asp Lys Asp Ala Pro Glu Pro Leu Asp  
115 120 125

Lys Val Ile Ile Leu Ser Pro Gly Ile Ser Asp Ala Thr Ala Pro Ala  
130 135 140

Trp Pro Pro Pro Gly Glu Asp Pro Gly Thr Thr Pro Pro Gly His Ser  
145 150 155 160

Val Pro Val Pro Ala Thr Glu Leu Gly Ser Thr Glu Leu Val Thr Thr  
165 170 175

Lys Thr Ala Gly Pro Glu Gln Gln  
180

<210> 17  
<211> 265  
<212> PRT  
<213> Homo sapiens

<400> 17

Met Ser Gly Leu Gly Arg Ser Arg Arg Gly Gly Arg Ser Arg Val Asp  
1 5 10 15

Gln Glu Glu Arg Phe Pro Gln Gly Leu Trp Thr Gly Val Ala Met Arg  
20 25 30

Ser Cys Pro Glu Glu Gln Tyr Trp Asp Pro Leu Leu Gly Thr Cys Met  
35 40 45

Ser Cys Lys Thr Ile Cys Asn His Gln Ser Gln Arg Thr Cys Ala Ala  
50 55 60

Phe Cys Arg Ser Leu Ser Cys Arg Lys Glu Gln Gly Lys Phe Tyr Asp  
65 70 75 80

His Leu Leu Arg Asp Cys Ile Ser Cys Ala Ser Ile Cys Gly Gln His  
85 90 95

Pro Lys Gln Cys Ala Tyr Phe Cys Glu Asn Lys Leu Arg Ser Pro Val  
100 105 110



Asn Leu Pro Pro Glu Leu Arg Arg Gln Arg Ser Gly Glu Val Glu Asn  
115 120 125

Asn Ser Asp Asn Ser Gly Arg Tyr Gln Gly Leu Glu His Arg Gly Ser  
130 135 140

Glu Ala Ser Pro Ala Leu Pro Gly Leu Lys Leu Ser Ala Asp Gln Val  
145 150 155 160

Ala Leu Val Tyr Ser Thr Leu Gly Leu Cys Leu Cys Ala Val Leu Cys  
165 170 175

Cys Phe Leu Val Ala Val Ala Cys Phe Leu Lys Lys Arg Gly Asp Pro  
180 185 190

Cys Ser Cys Gln Pro Arg Ser Arg Pro Arg Gln Ser Pro Ala Lys Ser  
195 200 205

Ser Gln Asp His Ala Met Glu Ala Gly Ser Pro Val Ser Thr Ser Pro  
210 215 220

Glu Pro Val Glu Thr Cys Ser Phe Cys Phe Pro Glu Cys Arg Ala Pro  
225 230 235 240

Thr Gln Glu Ser Ala Val Thr Pro Gly Thr Pro Asp Pro Thr Cys Ala  
245 250 255

Gly Arg Thr Ala Pro Pro Arg Glu Gly  
260 265